EXHIBIT

SEQ ID NO: 1

SEQ ID NO: 3

cataaaggac cacctacctg ggacgcgcag ttgggcggcg gactgggacg gcatgctgcg 60

g**t**gatgctgt cggtgatggt ctcttcctct ctggtcctga tcgtcttttt tctaggcgct 120

tccgaggagg cgaagccgg**c** **ga**cgacgacg acgataaaga atacaaagcc gcagtgtcgt 180

ccagaggatt acgcgaccag attgcaagat ctccgcgtca cctttcatcg agtaaaacct 240

*

acgt**tg**caac gtgaggacga ctactccgtg tggctcgacg gtacggtggt caaaggctgt 300

tggggatgca gcgtcatgga ctggttgttg aggcggtatc tggagatcgt gtt**c**cccgca 360

ggcgaccacg tctatcccgg actcaagacg gaattgcata gtatgcgctc gacgctagaa 420

tccatctaca aagacatgcg gcaatgcgta agtgtctctg tggcggcgct gtccgcacag 480

aggtaacaac gtgttcatag cacgctgttt tacttttgtc gggctcccag cctctgttag 540

gttgcggaga taagtccgtg attagtcggc tgtctcagga ggcggaaagg aaatcggata 600

acggcacgcg gaaaggtctc agcgagttgg acacgttgtt tagccgtctc gaagagtatc 660

tgcactcgag aaagtagcgt tgcgatttgc agtccgctcc ggtgtcgttc acccagttac 720

tttaataaac gtactgttta accrbmdcn 749

* 76 base pair insertion in Pestka sequence

SEQ 3: corresponds to residues 1-23 of SEQ ID NO: 1 as shown above.